Application No.: 10/568,206 3 Docket No.: 514572001600

## **AMENDMENTS TO THE CLAIMS**

1. (currently amended): A method for detecting a target <u>ribosomal ribonucleic</u> <del>nucleic</del> acid molecule (<u>rRNA</u>), said method <del>comprises</del> <u>comprising</u>:

- a) preparing a <u>bacterial</u> cell lysate comprising lysing a <u>bacterial</u> cell in a biological sample in a lysis buffer to release the target <u>nucleic acid rRNA</u> molecule from the <u>bacterial</u> cell;
- b) incubating the <u>bacterial</u> cell lysate from step a), without nucleic acid purification, with a <u>nucleic capture deoxyribonucleic</u> acid (<u>DNA</u>) probe immobilized on a solid substrate under conditions that allow <u>specific</u> hybridization between the target <u>nucleic acid rRNA</u> molecule and the <u>capture</u> probe, wherein the <u>nucleic acid capture</u> probe comprises a sequence complementary to the target <u>nucleic acid rRNA</u> molecule;
- c) assessing hybridization between the target nucleic acid rRNA molecule and the capture DNA probe to determine the presence, absence and/or amount of the target nucleic acid rRNA molecule,

wherein the hybridization between the target rRNA molecule and the capture probe is assessed by determining specific binding of a reporter to the target rRNA molecule, wherein the reporter comprises a reporter DNA probe complementary to the target rRNA molecule and a detectable marker selected from the group consisting of a fluorescein, an isotope, a biotin, a digoxin, a gold colloid, a magnetic bead, a electrochemical label, and a chemiluminescent label; and steps a) through c) can be completed in 90 minutes or less.

- 2. (currently amended): The method of claim 1, wherein the <u>bacterial</u> cell is lysed in the lysis buffer by a physical method.
- 3. (original): The method of claim 2, wherein the physical method is selected from the group consisting of grinding, ultrasonic lysing, lysing with high temperature, and freezing.
- 4. (currently amended): The method of claim 1, wherein the <u>bacterial</u> cell is lysed in the lysis buffer by a chemical method.

5. (original): The method of claim 4, wherein the chemical method is lysing with a protein denaturant or a detergent.

- 6. (currently amended): The method of claim 1, wherein the <u>bacterial</u> cell is lysed in the lysis buffer by a biological method.
- 7. (original): The method of claim 6, wherein the biological method is lysing with a proteinase or a lysozyme.
- 8. (currently amended): The method of claim 1, wherein the <u>bacterial</u> cell is lysed by any combination of a physical <u>method</u>, a chemical <u>method</u>, and a biological method.
- 9. (currently amended): The method of claim 1, wherein the cell lysate is incubated with the capture probe immobilized on the substrate in the lysis buffer for hybridization.
- 10. (currently amended): The method of claim 1, wherein an agent that aids for hybridization is added to the cell lysate before the cell lysate is incubated with the <u>capture</u> probe.
- 11. (currently amended): The method of claim 10, wherein the agent is selected from the group consisting of NaCl sodium chloride, eitrate sodium citrate, and SDS sodium dodecyl sulfate.
- 12. (currently amended): The method of claim 1, wherein the biological sample is a sample selected from the group consisting of a non-virus biological organism, a biological tissue, a eukaryotic cell, and a prokaryotic cell.

## 13. (canceled)

14. (original): The method of claim 1, wherein the solid substrate comprises a material selected from the group consisting of a nylon film, a pyroxylin film, a silicon, a glass, a ceramic, a metal, a plastic, and a combination thereof.

15. (currently amended): The method of claim 1, wherein the solid substrate comprises a plurality of nucleic acid capture probes, and wherein the plurality of the nucleic acid capture probes are immobilized on the solid substrate to form an array.

- 16. (currently amended): The method of claim 15, wherein the plurality of the nucleic acid capture probes have different nucleotide sequences.
- 17. (currently amended): The method of claim 16, wherein the number of different capture probes is from about 2 to about 100,000.
- 18. (currently amended): The method of claim 15, wherein the array [[is]] <u>has an area ranging</u> from about 0.01 mm<sup>2</sup> to about 100 cm<sup>2</sup>.
- 19. (currently amended)): The method of claim 15, wherein the array is selected from the group consisting of a two-dimensional array[[,]] and a three-dimensional array, and a four dimensional array.
- 20. (currently amended): The method of claim 1, wherein the nucleic acid capture probe immobilized on the solid substrate comprises a single-stranded oligonucleotide or a double-stranded PCR product.
- 21. (currently amended): The method of claim 1, wherein the <u>bacterial</u> cell lysate comprises an agent selected from the group consisting of a detergent, a protein denaturant, a buffer, a nuclease inhibitor, a salt, and a combination thereof.
  - 22. (canceled)
- 23. (new): The method of claim 1, wherein the reporter is added to the bacterial cell lysate before the bacterial cell lysate has been incubated with the capture probe.

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24. (new): The method of claim 1, wherein the reporter is added to the bacterial cell lysate after bacterial the cell lysate has been incubated with the capture probe.